DT09 Rec'd PCT/PT0 3 0 NOV 2004

SEQUENCE LISTING

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- <130> D3-A0203P
- <150> JP 2002-161964
- <151> 2002-06-03
- <160> 63
- <170> PatentIn version 3.1
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- <211> 10
- <212> DNA
- <213> Sendai virus
- <400> 1

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115 ---

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Tyr	Thr	Leu	Ser	Ser	Ser	Val	Thr	Val	Pro	Ser	Ser	Thr	Trp	Pro	Ser	
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Ser	Val	Val	Cys		Leu	Asn	Asn	Phe		Pro	Lys	Asp	lle		Val	
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415 420 425

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1397

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430

435

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455

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Thr Phe Thr Asn Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His
50 55 60

Gly Leu Glu Trp Ile Gly Asp Ile Tyr Pro Gly Gly Gly Tyr Thr Asn
65 70 75 80

Tyr Asn Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Thr Ser

85 90 95

Ser Ser Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser

100 105 110

Ala Val Tyr Phe Cys Ala Arg Phe Tyr Tyr Gly Ser Ser Tyr Trp Tyr

115
120 --125

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Thr Asn Ser Met Val Thr Leu Gly Cys Leu Val Lys Gly Tyr Phe Pro 165 170 175

Glu Pro Val Thr Val Thr Trp Asn Ser Gly Ser Leu Ser Ser Gly Val
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His Thr Phe Pro Ala Val Leu Gln Ser Asp Leu Tyr Thr Leu Ser Ser 195 200 205

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90

95

Ser Ser Leu His Pro Asp Asp Val Ala Thr Tyr Tyr Cys Gln Asn Val

100 105 110

Leu Ser Thr Pro Arg Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys

115 120 125

Arg Ala Asp Ala Ala Pro Thr Val Ser Ile Phe Pro Pro Ser Ser Glu
130 135 140

Gln Leu Thr Ser Gly Gly Ala Ser Val Val Cys Phe Leu Asn Asn Phe 145 150 155 160

Tyr Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg
165 170 175

Gln Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser

180 185 190

Thr Tyr Ser Met Ser Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu

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cctggad	eatg 70
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35 40 45

Lys Leu Leu Ile Phe Ala Ala Ser Asn Val Glu Ser Gly Val Pro Ala 50 55 60

Arg Phe Ser Gly Ser Gly Ser Gly Thr Asn Phe Ser Leu Asn Ile His

70

75

80

Pro Val Asp Glu Asp Asp Val Ala Met Tyr Phe Cys Gln Gln Ser Arg

85 90 95

Lys Val Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg

100 105 110

Gly	Gly	Gly	Gly	Ser	Gly	Gly	Gly	Gly	Ser	Gly	Gly	Gly	Gly	Ser	Gln
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Val Lys Leu Gln Gln Ser Gly Pro Gly Leu Val Thr Pro Ser Gln Ser 130 135 140

Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu Ser Asp Tyr Gly
145 150 155 160

Val His Trp Val Arg Gln Ser Pro Gly Gln Gly Leu Glu Trp Leu Gly
165 170 175

Val Ile Trp Ala Gly Gly Gly Thr Asn Tyr Asn Ser Ala Leu Met Ser 180 185 190

Arg Lys Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu Lys
195 200 205

Met Asn Ser Leu Gln Ala Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg
210 215 220

Asp Lys Gly Tyr Ser Tyr Tyr Tyr Ser Met Asp Tyr Trp Gly Gln Gly
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Thr Thr Val Thr Val Ser Ser

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cctcag	tcat aatgtccaga ggatctagac cgtagtaaga aaaacttagg gtgaaagttc 12	20
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